

99274Cseq
SEQUENCE LISTING

<110> Boutand, Ariel

<120> Recombinant Laminin 5

<130> 99-274-C

<140> 09/560,385

<141> 2000-04-28

<160> 36

<170> PatentIn Ver. 2.0

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 Page 13

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gaa Glu	aat Asn 930	gtt Val	ctg Leu	agc Ser	ttg Leu	tac Tyr 935	aac Asn	ttc Phe	aaa Lys	aaa Lys	aca Thr 940	ttc Phe	aat Asn	ctc Leu	aac Asn	2832
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cca Pro 1025	aaa Lys	gag Glu	aga Arg	gga Gly	gtt Val 1030	gga Gly	gac Asp	gcc Ala	ata Ile 1035	aac Asn	aac Asn	ggc Gly	aga Arg	gac Asp	cat His 1040	3120
tcg Pro	att Tyr	cag Glu	atc Ile	aaa Lys	att Tyr	gga Glu	aaa Lys	ctc Leu	caa Gln	aag Lys	cgt Arg	atg Met	tgg Val	ata Ile	aat Asn	3168

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Thr Tyr Tyr Leu Gly Gly Ile Pro Ile Ala Ile Arg Glu Arg Phe Asn	
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Gln Thr Thr Ser Ser Arg Gly Leu Val Phe His Thr Gly Thr Lys Asn	
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Pro His Thr Asn Ser Phe Ala Thr Gly Cys Val Val Asn Gly Gly Asp
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Cys Ala Pro Gly Tyr Phe Gly Asn Pro Gln Lys Phe Gly Gly Ser Cys
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Gln Pro Cys Ser Cys Asn Ser Asn Gly Gln Leu Gly Ser Cys His Pro
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Leu Thr Gly Asp Cys Ile Asn Gln Glu Pro Lys Asp Ser Ser Pro Ala
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Gly Leu Ser Ala Ser Ala Gly Leu Leu Glu Gln Met Arg His Met Glu
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Asp Leu Asn Gln Glu Phe Glu Thr Leu Gln Glu Lys Ala Gln Val Asn
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Ser Arg Lys Ala Gln Thr Leu Asn Asn Asn Val Asn Arg Ala Thr Gln
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Cys Asn Gly His Ser Asn Gln Cys Gln Asp Gly Ser Gly Ile Cys Val
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Asn	Gly	His	Ser	Asn	Arg	Cys	Gln	Asp	Gly	Ser	Gly	Val	Cys	Ile	Asn	95	
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tgt	cag	cac	aac	aca	gct	ggg	gag	cac	tgt	gag	cgt	tgc	aag	agg	ggt	394	
Cys	Gln	His	Asn	Thr	Ala	Gly	Glu	His	Cys	Glu	Arg	Cys	Lys	Arg	Gly	110	
			100					105									
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Tyr	Tyr	Gly	Ser	Ala	Ile	His	Gly	Ser	Cys	Arg	Val	Cys	Pro	Cys	Pro	125	
		115					120					125					
cac	acc	aac	agc	ttt	gcc	act	ggc	tgt	gct	gtg	gat	gga	gga	gct	gtg	490	
His	Thr	Asn	Ser	Phe	Ala	Thr	Gly	Cys	Ala	Val	Asp	Gly	Gly	Ala	Val	140	
	130					135											
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Arg	Cys	Ala	Cys	Lys	Pro	Gly	Tyr	Thr	Gly	Ala	Gln	Cys	Glu	Arg	Cys	160	
145					150				155								
gca	cca	gga	tat	ttt	ggg	aac	ccc	cag	aaa	ttt	gga	ggt	agc	tgc	caa	586	
Ala	Pro	Gly	Tyr	Phe	Gly	Asn	Pro	Gln	Lys	Phe	Gly	Gly	Ser	Cys	Gln	175	
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cca	tgc	aat	tgc	aac	agt	aat	ggc	cag	ttt	ggc	act	tgt	gat	ccc	cta	634	
Pro	Cys	Asn	Cys	Asn	Ser	Asn	Gly	Gln	Phe	Gly	Thr	Cys	Asp	Pro	Leu	180	
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Thr	Gly	Asp	Cys	Val	Ser	Gln	Glu	Pro	Lys	Asp	Gly	Ser	Pro	Ala	Glu	195	
		195					200					205					
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Glu	Cys	Asp	Asp	Cys	Asp	Ser	Cys	Val	Met	Thr	Leu	Leu	Asn	Asp	Leu	210	
	210					215					220						
gtc	ccc	atg	ggt	gag	gaa	ctc	gcc	ctg	gtg	aaa	tca	aaa	ctt	cag	ggg	778	
Val	Pro	Met	Gly	Glu	Glu	Leu	Ala	Leu	Val	Lys	Ser	Lys	Leu	Gln	Gly	225	
225				230					235						240		
ctg	agt	gtg	aac	act	ggt	tct	ctg	gaa	cag	atc	cgg	cat	gtg	gag	atg	826	
Leu	Ser	Val	Asn	Thr	Gly	Ser	Leu	Glu	Gln	Ile	Arg	His	Val	Glu	Met	245	
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cag	gcc	aag	gac	ctg	agg	aac	cag	ctg	ctt	ggc	ttc	cgt	tcc	gcc	atc	874	
Gln	Ala	Lys	Asp	Leu	Arg	Asn	Gln	Leu	Leu	Gly	Phe	Arg	Ser	Ala	Ile	260	
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tcc	agt	cac	ggg	tcc	caa	atg	gac	ggc	ctg	gaa	aaa	gaa	ctc	agt	cat	922	
Ser	Ser	His	Gly	Ser	Gln	Met	Asp	Gly	Leu	Glu	Lys	Glu	Leu	Ser	His	275	
		275					280					285					
ttg	tac	cag	gaa	ttc	gaa	act	ttg	caa	gaa	aag	gcg	cag	gtc	aat	tcc	970	
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gcc	aaa	gag	ttg	gac	atg	aag	att	aaa	aac	ata	ctt	acg	aat	gtg	cac	1066
Ala	Lys	Glu	Leu	Asp	Met	Lys	Ile	Lys	Asn	Ile	Leu	Thr	Asn	Val	His	
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att	ctc	ctg	aag	cag	atc	gct	cgg	cca	ggc	gga	gaa	gga	atg	gac	ttg	1114
Ile	Leu	Leu	Lys	Gln	Ile	Ala	Arg	Pro	Gly	Gly	Glu	Gly	Met	Asp	Leu	
			340					345					350			
ccg	gtg	ggc	gac	tgg	tcc	agg	gag	tcg	gcg	gaa	gct	cag	cgc	atg	ttg	1162
Pro	Val	Gly	Asp	Trp	Ser	Arg	Glu	Ser	Ala	Glu	Ala	Gln	Arg	Met	Leu	
		355					360					365				
cgg	gag	ctg	cga	ggc	cga	gac	ttt	aaa	aag	cac	ctc	caa	gaa	gca	gag	1210
Arg	Glu	Leu	Arg	Gly	Arg	Asp	Phe	Lys	Lys	His	Leu	Gln	Glu	Ala	Glu	
	370					375					380					
gcc	cag	aaa	atg	gaa	gcc	cag	ctc	tta	ctg	aac	cga	atc	agg	acc	tgg	1258
Ala	Gln	Lys	Met	Glu	Ala	Gln	Leu	Leu	Leu	Asn	Arg	Ile	Arg	Thr	Trp	
385					390					395					400	
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Leu	Glu	Ser	His	Gln	Val	Glu	Asn	Asn	Gly	Leu	Leu	Lys	Asn	Ile	Arg	
			405						410					415		
gat	tca	tta	aat	gat	tat	gaa	gcc	aaa	ctt	cag	gac	ctg	cgt	tcc	gtg	1354
Asp	Ser	Leu	Asn	Asp	Tyr	Glu	Ala	Lys	Leu	Gln	Asp	Leu	Arg	Ser	Val	
			420					425					430			
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Leu	Gln	Glu	Ala	Ala	Ala	Gln	Gly	Lys	Gln	Ala	Thr	Gly	Leu	Asn	His	
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gaa	aat	gag	ggg	gtc	cta	gga	gcc	atc	cag	aga	caa	atg	aag	gaa	atg	1450
Glu	Asn	Glu	Gly	Val	Leu	Gly	Ala	Ile	Gln	Arg	Gln	Met	Lys	Glu	Met	
	450					455					460					
gat	tcc	ctg	aag	aag	tac	ctc	acc	gag	cac	ctg	gcc	aca	gca	gac	gct	1498
Asp	Ser	Leu	Lys	Lys	Tyr	Leu	Thr	Glu	His	Leu	Ala	Thr	Ala	Asp	Ala	
465					470					475					480	
tcc	ctg	ctg	caa	acc	aac	agt	cta	ctg	cag	cgg	atg	gac	acg	agc	cag	1546
Ser	Leu	Leu	Gln	Thr	Asn	Ser	Leu	Leu	Gln	Arg	Met	Asp	Thr	Ser	Gln	
			485						490					495		
aag	gag	tat	gaa	agc	tta	gct	gct	gct	tta	aac	gga	gca	aga	cag	gaa	1594
Lys	Glu	Tyr	Glu	Ser	Leu	Ala	Ala	Ala	Leu	Asn	Gly	Ala	Arg	Gln	Glu	
			500					505					510			
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Leu	Asn	Asp	Gln	Val	Arg	Glu	Leu	Ser	Arg	Ser	Gly	Gly	Lys	Ala	Pro	
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Leu	Val	Ala	Glu	Ala	Glu	Lys	His	Ala	Gln	Ser	Leu	Gln	Glu	Leu	Ala	
	530					535					540					
aag	cag	ctg	gaa	gag	ata	aag	aga	aac	acc	agt	ggg	gat	gag	tcg	gtg	1738
Lys	Gln	Leu	Glu	Glu	Ile	Lys	Arg	Asn	Thr	Ser	Gly	Asp	Glu	Ser	Val	
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Ile	Arg	Ala	Ala	Glu	Asp	Ala	Ala	Gly	Lys	Ala	Asp	Ser	Ala	Ser	Glu	
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Ser	Ala	Phe	Gln	Thr	Val	Ile	Lys	Glu	Asp	Leu	Pro	Arg	Arg	Ala	Lys	
		595					600					605				
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Thr	Leu	Ser	Ser	Asp	Ser	Glu	Glu	Leu	Leu	Asn	Glu	Ala	Lys	Met	Thr	
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Arg	Lys	Arg	Leu	Gln	Gln	Glu	Ile	Asn	Pro	Ala	Leu	Asn	Ser	Leu	Gln	
625					630					635					640	
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Gln	Thr	Leu	Lys	Thr	Val	Ser	Val	Gln	Lys	Asp	Leu	Leu	Asp	Ala	Asn	
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gtc	act	gct	gtc	cgt	aat	gac	ctt	cgt	ggg	atc	cag	aga	ggt	gat	att	2074
Val	Thr	Ala	Val	Arg	Asn	Asp	Leu	Arg	Gly	Ile	Gln	Arg	Gly	Asp	Ile	
			660					665					670			
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Asp	Ser	Val	Val	Ser	Gly	Ala	Lys	Ser	Met	Val	Arg	Lys	Ala	Asn	Gly	
		675					680					685				
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Ile	Thr	Ser	Glu	Val	Leu	Asp	Gly	Leu	Ser	Pro	Ile	Gln	Thr	Asp	Leu	
	690					695					700					
gga	agg	att	aag	gac	agc	tac	ggg	agc	aca	cgg	cat	gag	gac	ttc	aac	2218
Gly	Arg	Ile	Lys	Asp	Ser	Tyr	Gly	Ser	Thr	Arg	His	Glu	Asp	Phe	Asn	
705					710					715					720	
aaa	gct	ctg	att	gac	gcc	aat	aac	tca	gta	aag	aaa	tta	acc	aag	aag	2266
Lys	Ala	Leu	Ile	Asp	Ala	Asn	Asn	Ser	Val	Lys	Lys	Leu	Thr	Lys	Lys	
				725					730					735		
ttg	cct	gat	ctt	ttt	gtc	aag	att	gaa	agc	atc	aat	caa	cag	ttg	ctg	2314
Leu	Pro	Asp	Leu	Phe	Val	Lys	Ile	Glu	Ser	Ile	Asn	Gln	Gln	Leu	Leu	
			740					745					750			
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Pro	Leu	Gly	Asn	Ile	Ser	Asp	Asn	Val	Asp	Arg	Ile	Arg	Glu	Leu	Ile	
		755					760					765				
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Thr	Gln	Ala	Arg	Asp	Ala	Ala	Asn	Lys	Val	Ala	Ile	Pro	Met	Arg	Phe	
	770				775						780					
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Asn	Gly	Lys	Ser	Gly	Val	Glu	Val	Arg	Leu	Pro	Asn	Asp	Leu	Glu	Asp	
785				790						795					800	
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Leu	Lys	Gly	Tyr	Thr	Ser	Leu	Ser	Leu	Phe	Leu	Gln	Arg	Pro	Asp	Leu	

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805				810				815								
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Arg	Glu	Asn	Gly	Gly	Thr	Glu	Asp	Met	Phe	Val	Met	Tyr	Leu	Gly	Asn	
			820					825					830			
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Lys	Asp	Ala	Ser	Lys	Asp	Tyr	Ile	Gly	Met	Ala	Val	Val	Asp	Gly	Gln	
		835					840					845				
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Leu	Thr	Cys	Val	Tyr	Asn	Leu	Gly	Asp	Arg	Glu	Ala	Glu	Val	Gln	Ile	
	850					855					860					
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Asp	Gln	Val	Leu	Thr	Glu	Ser	Glu	Ser	Gln	Glu	Ala	Val	Met	Asp	Arg	
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Val	Lys	Phe	Gln	Arg	Ile	Tyr	Gln	Phe	Ala	Lys	Leu	Asn	Tyr	Thr	Lys	
			885						890					895		
gaa	gcc	acg	tcc	aat	aaa	ccc	aaa	gct	ccc	gcg	gtc	tac	gac	ctg	gag	2794
Glu	Ala	Thr	Ser	Asn	Lys	Pro	Lys	Ala	Pro	Ala	Val	Tyr	Asp	Leu	Glu	
			900					905					910			
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Gly	Gly	Ser	Ser	Asn	Thr	Leu	Leu	Asn	Leu	Asp	Pro	Glu	Asp	Ala	Val	
		915				920						925				
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Phe	Tyr	Val	Gly	Gly	Tyr	Pro	Pro	Asp	Phe	Glu	Leu	Pro	Ser	Arg	Leu	
	930					935					940					
cgg	ttc	cct	cca	tac	aaa	ggc	tgt	atc	gaa	cta	gat	gac	ctc	aat	gaa	2938
Arg	Phe	Pro	Pro	Tyr	Lys	Gly	Cys	Ile	Glu	Leu	Asp	Asp	Leu	Asn	Glu	
945					950					955					960	
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Asn	Val	Leu	Ser	Leu	Tyr	Asn	Phe	Lys	Thr	Thr	Phe	Asn	Leu	Asn	Thr	
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			980					985					990			
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		995				1000						1005				
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Pro	Phe	Pro	Asn	Phe	Ile	Gln	Thr	Ile	Gln	Thr	Thr	Val	Asp	Arg	Gly	
	1010				1015					1020						
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Leu	Leu	Phe	Phe	Ala	Glu	Asn	Gln	Asp	Asn	Phe	Ile	Ser	Leu	Asn	Ile	
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Ile	Leu	Ile	Thr	Ile	Gly	Lys	Leu	Gln	Lys	Arg	Met	Trp	Ile	Asn	Val		
			1075				1080					1085					
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Gly	Val	Val	Arg	Leu	Asn	Asp	Thr	Val	Gly	Val	Thr	Lys	Lys	Cys	Ser		
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			1155				1160					1165					
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Thr	Tyr	Met	Asp	Gly	Leu	Leu	His	His	Val	Ser	Val	Ile	Ser	Asp	Thr		
			1235			1240						1245					
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			1250			1255					1260						
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			1265			1270				1275				1280			
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Gln	Ser	Pro	Glu	Val	Leu	Asp	Leu	Ala	Ser	Lys	Ser	Thr	Lys	Lys	Asp		
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Ala Ser Leu Gly Gly Cys Ser Leu Asn Lys Pro Pro Phe Leu Met Leu	
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Phe Lys Ser Pro Lys Arg Phe Asn Lys Gly Arg Ile Phe Asn Val Asn	
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Gln Leu Met Gln Asp Ala Pro Gln Ala Thr Arg Ser Thr Glu Ala Trp	
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Gln Asp Gly Arg Ser Cys Leu Pro Pro Leu Asn Thr Lys Ala Ser His	
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Arg Ala Leu Gln Phe Gly Asp Ser Pro Thr Ser His Leu Leu Lys	
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Leu Pro Gln Glu Leu Leu Lys Pro Arg Ser Gln Phe Ser Leu Asp Ile	
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Gln Thr Thr Ser Pro Lys Gly Leu Val Phe Tyr Ala Gly Thr Lys Asp	
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Gly Ala Gly Gly Lys Lys Leu Arg Leu Arg Ser Lys Glu Arg Tyr His	
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Asn Cys Gln 65	His Asn Thr 70	Ala Gly Glu His Cys 75	Glu Arg Cys Lys Arg 80
Gly Tyr Tyr 85	Gly Ser Ala Ile His 90	Cys Arg Val Cys Pro 95	Cys
Pro His Thr 100	Asn Ser Phe Ala Thr 105	Gly Cys Ala Val Asp 110	Gly Gly Ala
Val Arg Cys 115	Ala Cys Lys Pro Gly 120	Tyr Thr Gly Ala Gln 125	Cys Glu Arg
Cys Ala Pro 130	Gly Tyr Phe Gly 135	Asn Pro Gln Lys Phe 140	Gly Gly Ser Cys
Gln Pro Cys 145	Asn Cys Asn Ser 150	Asn Gly Gln Phe 155	Gly Thr Cys Asp Pro 160
Leu Thr Gly 165	Asp Cys Val Ser 170	Gln Glu Pro Lys 175	Asp Gly Ser Pro Ala
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Gly Leu Ser 210	Val Asn Thr Gly 215	Ser Leu Glu Gln Ile 220	Arg His Val Glu
Met Gln Ala 225	Lys Asp Leu Arg 230	Asn Gln Leu Leu 235	Gly Phe Arg Ser Ala 240
Ile Ser Ser 245	His Gly Ser Gln 250	Met Asp Gly Leu 255	Glu Lys Glu Leu Ser
His Leu Tyr 260	Gln Glu Phe Glu 265	Thr Leu Gln Glu 270	Lys Ala Gln Val Asn
Ser Arg Lys 275	Ala Gln Thr Leu 280	Tyr Asn Asn Ile 285	Asp Thr Thr Ile Gln
Asn Ala Lys 290	Glu Leu Asp Met 295	Lys Ile Lys Asn 300	Ile Leu Thr Asn Val
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 Arg Leu Gly Gln Ser Ser Met Leu Gly Glu Gln Gly Ala Arg Ile Gln
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Cys Lys Cys Asp Ser Arg Gln Pro His Asn Tyr Tyr Ser His Arg Val
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Arg	Pro	Asn	Ala	Arg	Leu	Asn	Gly	Gly	Lys	Val	Gln	Leu	Asn	Leu	Met	195	200	205	
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Gly	Gln	Asp	Ala	His	Glu	Cys	Gln	Arg	Cys	Asp	Cys	Asn	Gly	His	Ser	325	330	335	
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99274Cseq

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 Gln Cys Pro₅₁₅ Cys Arg Glu Gly Phe₅₂₀ Gly Gly Leu Met Cys₅₂₅ Ser Ala Ala
 Ala Ile₅₃₀ Arg Gln Cys Pro Asp₅₃₅ Arg Thr Tyr Gly Asp₅₄₀ Val Ala Thr Gly
 Cys₅₄₅ Arg Ala Cys Asp Cys₅₅₀ Asp Phe Arg Gly Thr₅₅₅ Glu Gly Pro Gly Cys₅₆₀
 Asp Lys Ala Ser Gly₅₆₅ Arg Cys Leu Cys Arg₅₇₀ Pro Gly Leu Thr Gly₅₇₅ Pro
 Arg Cys Asp Gln₅₈₀ Cys Gln Arg Gly Tyr₅₈₅ Cys Asn Arg Tyr Pro₅₉₀ Val Cys
 Val Ala Cys₅₉₅ His Pro Cys Phe Gln₆₀₀ Thr Tyr Asp Ala Asp₆₀₅ Leu Arg Glu
 Gln Ala₆₁₀ Leu Arg Phe Gly Arg₆₁₅ Leu Arg Asn Ala Thr₆₂₀ Ala Ser Leu Trp
 Ser Gly Pro Gly Leu Glu₆₃₀ Asp Arg Gly Leu Ala₆₃₅ Ser Arg Ile Leu Asp₆₄₀
 Ala Lys Ser Lys Ile₆₄₅ Glu Gln Ile Arg Ala₆₅₀ Val Leu Ser Ser Pro₆₅₅ Ala
 Val Thr Glu Gln₆₆₀ Glu Val Ala Gln Val₆₆₅ Ala Ser Ala Ile Leu₆₇₀ Ser Leu
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 Glu Gln Ser Ala₇₄₀ Gln Ala Ala Gln Gln₇₄₅ Val Ser Asp Ser Ser₇₅₀ Arg Leu
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Gln Met Ile Arg Ala Ala Glu Glu Ser Ala Ser Gln Ile Gln Ser Ser
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Glu Asp Val Arg Arg Thr Arg Leu Leu Ile Gln Gln Val Arg Asp Phe
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Ala Val Leu Ala Leu Trp Leu Pro Thr Asp Ser Ala Thr Val Leu Gln
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Lys Met Asn Glu Ile Gln Ala Ile Ala Ala Arg Leu Pro Asn Val Asp
945 950 955 960

Leu Val Leu Ser Gln Thr Lys Gln Asp Ile Ala Arg Ala Arg Arg Leu
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Gln Ala Glu Ala Glu Glu Ala Arg Ser Arg Ala His Ala Val Glu Gly
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Lys Leu Val Thr Ser Met Thr Lys Gln Leu Gly Asp Phe Trp Thr Arg
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Ser Val Lys Thr Glu Ala Glu Glu Leu Phe Gly Glu Thr Met Glu Met
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145 150 155

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Pro	Asn	Ala	Arg	Leu	Asn	Gly	Gly	Lys	Val	Gln	Leu	Asn	Leu	Met	Asp	
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Ser	Ala	Gln	Arg	Leu	Glu	Thr	Gln	Val	Ser	Ala	Ser	Arg	Ser	Gln	Met	
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 Cys Cys Lys Cys Asp Ser Arg Gln Pro His Asn Tyr Tyr Ser His Arg
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 Phe Gln Leu Gln Glu Val Met Met Glu Phe Gln Gly Pro Met Pro Ala
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 Met Asp Leu Val Ser Gly Ile Pro Ala Thr Gln Ser Gln Lys Ile Gln
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610

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<400> 26

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 Ser Ser Ala Gln Arg Leu Asp Pro Val Tyr Phe Val Ala Pro Ala Lys
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Phe	Leu	Gly	Asn ₂₆₀	Gln	Gln	Val	Ser	Tyr ₂₆₅	Gly	Gln	Ser	Leu	Ser ₂₇₀	Phe	Asp
Tyr	Arg	Val ₂₇₅	Asp	Arg	Gly	Gly	Arg ₂₈₀	Gln	Pro	Ser	Ala	Tyr ₂₈₅	Asp	Val	Ile
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 Tyr Ser Gly Asp Glu Asn Pro Asp Ile Glu Cys Ala Asp Cys Pro Ile
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 Cys His Asn Gly Phe Ser Cys Ser Val Met Pro Glu Thr Glu Glu Val
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99274Cseq
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Cys	Tyr	Asn 595	Gln	Val	Lys	Ile	Gln 600	Met	Asp	Gln	Phe	Thr 605	Gln	Gln	Leu		
cag	agc	ctg	gag	gcc	ctg	gtt	tca	aag	gct	cag	ggt	ggt	ggt	ggt	ggt	1872	
Gln	Ser 610	Leu	Glu	Ala	Leu	Val 615	Ser	Lys	Ala	Gln	Gly 620	Gly	Gly	Gly	Gly		
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Gly 625	Thr	Val	Pro	Val	Gln 630	Leu	Glu	Gly	Arg	Ile 635	Glu	Gln	Ala	Glu	Gln 640		
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Tyr	Lys	Thr 675	Arg	Leu	Asp	Asp	Leu 680	Lys	Met	Thr	Ala	Glu 685	Arg	Ile	Arg		
gcc	ctg	ggc	agt	cag	cat	cag	aac	aga	gtt	cag	gat	acg	agc	aga	ctc	2112	
Ala	Leu 690	Gly	Ser	Gln	His	Gln 695	Asn	Arg	Val	Gln	Asp 700	Thr	Ser	Arg	Leu		
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Asn	Thr	Asn	Ile	His 725	Ser	Ser	Glu	His	Tyr 730	Val	Gly	Pro	Asn	Asp 735	Phe		
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Lys	Ser	Leu	Ala 740	Gln	Glu	Ala	Thr	Arg 745	Lys	Ala	Asp	Ser	His 750	Ala	Glu		
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Ser	Ala	Asn 755	Ala	Met	Lys	Gln	Leu 760	Ala	Arg	Glu	Thr	Glu 765	Asp	Tyr	Ser		
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Ser 785	Gly	Ser	Trp	Asp	Ser 790	Ser	Val	Val	Gln	Gly 795	Leu	Met	Gly	Lys	Leu 800		
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 35 40 45
 Glu Gly Phe Tyr Gln His Gln Ser Lys Ser Arg Cys Leu Pro Cys Asn
 50 55 60
 Cys His Ser Lys Gly Ser Leu Ser Ala Gly Cys Asp Asn Ser Gly Gln
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 Cys Arg Cys Lys Pro Gly Val Thr Gly Gln Arg Cys Asp Gln Cys Gln
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 Pro Gly Phe His Met Leu Thr Asp Ala Gly Cys Thr Arg Asp Gln Gly
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 Gln Leu Asp Ser Lys Cys Asp Cys Asp Pro Ala Gly Ile Ser Gly Pro
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 Glu Gly Cys Thr Gln Cys Phe Cys Tyr Gly His Ser Ala Ser Cys His
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 Pro His Asp Pro Arg Ser Cys Lys Pro Cys Pro Cys His Asn Gly Phe
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 Pro Pro Gly Val Thr Gly Ala Arg Cys Glu Leu Cys Ala Asp Gly Phe
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 Tyr Lys Thr Arg Leu Asp Asp Leu Lys Met Thr Ala Glu Arg Ile Arg
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 705 710 715 720
 Asn Thr Asn Ile His Ser Ser Glu His Tyr Val Gly Pro Asn Asp Phe
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Lys Val Ala Asp Ala Ser Asp Lys Thr Gln Gln Ala Glu Thr Ala Leu
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